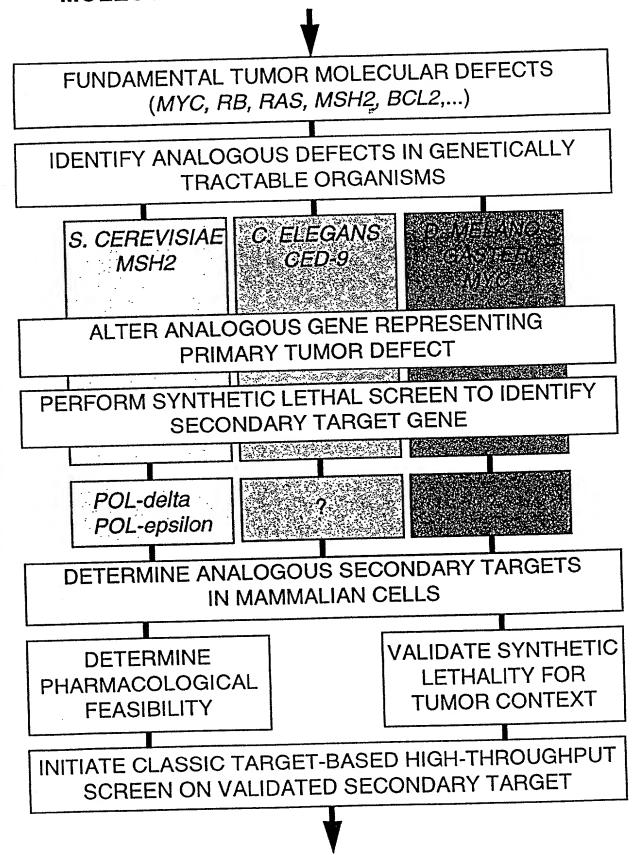
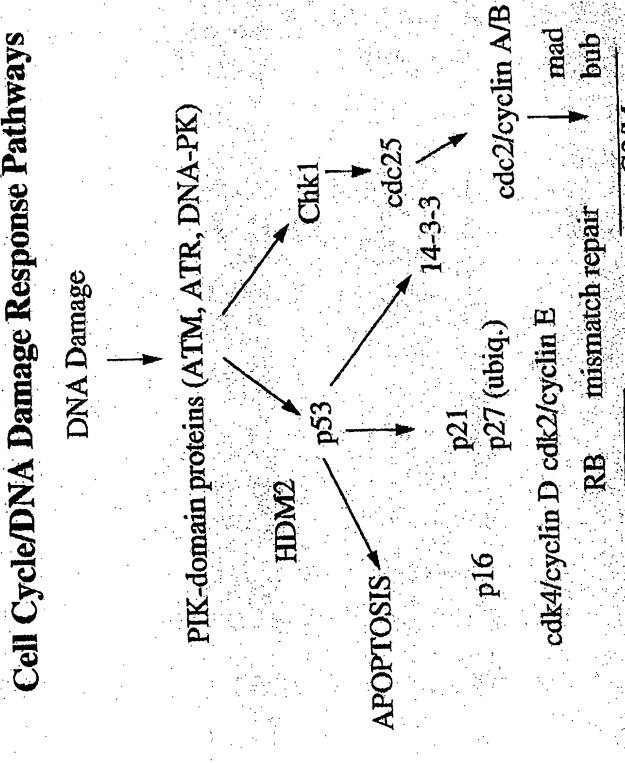
MOLECULAR ALTERATIONS IN TUMORS

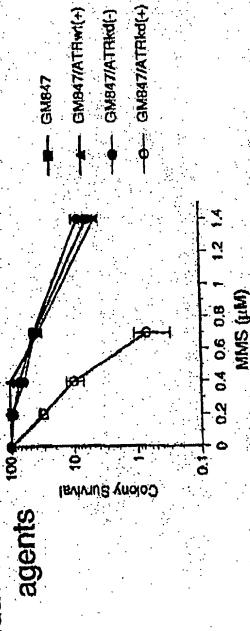


ANTI-CANCER DRUGS BASED ON TUMOR CONTEXT



MAMMALIAN CELL EVALUATION OF ATR AS A TARGET

Overexpression of ATR-KD not tolerated in human tumor cell lines (MCF-7, A549) 2. Inducible ATR-KD sensitizes cells to DNA damaging



3. LCK promoter driven ATR-KD transgenic mice have cells stably expressing ATR-KD in thymus

Synthetic lethality:

• Use primary defect as a selective context to kill tumor cells with an alteration in gene A.

Combined defects in gene A and gene B kill tumor cells while disrupting gene B activity alone has no effect on normal cells.

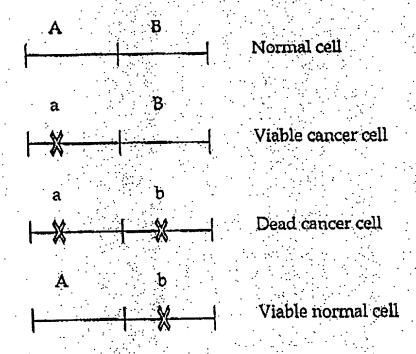


Figure 5

Human genes altered in tumors and their relatives in model genetic systems. Genes that are not structural homologs but act in analogous pathways (such as human p53 and S. cerevisiae RAD9) are shown in brackets. Saccharomyces cerevisiae genes are designated with superscript Sc, S. pombe with Sp, C. elegans with Ce, and D. melanogaster with Dm. Because of space limitations, this is only a representative list of genes mutated in tumors that have genetic analogs in model systems.

Function	Human genes	Model system analogs: structural homologs or related biological roles
DNA damage checkpoint	p53	[RAD9 ^{Sc} , rad1+ ^{Sp}]
# # # # # # # # # # # # # # # # # # #	ATM	MEC1 ^{Sc} , TEL1 ^{Sc} , rad3 ^{3+Sp} , mei-41 ^{Dm}
DNA mismatch	MSH2, MLH1	MSH2 ^{Sc} , MLH1 ^{Sc}
Nucleotide excision repair	XP-A, XP-B	RAD14 ^{sc} , RAD25 ^{sc}
Of methylguanine reversal	MGMT	MGT1 ^{Sc}
Double-strand break repair	BRCA2, BRCA1	[RAD51 ^{Sc} , RAD54 ^{Sc}]
DNA helicase	BLM	SGS1Sc, rgh1+So
Growth factor	RAS	RAS1Sc, RAS2Sc, let-60Ce
signaling		
	NF1	IRA1 ^{Sc} , IRA2 ^{Sc}
	MYC	dMyc ^{Dm}
0 -4	· PTH	patched ^{Om}
Cell cycle control	Cyclin D, Cyclin E	CLN1 ^{Sc} , CLN2 ^{Sc} , Cyclin D ^{Rm} , Cyclin E ^{Dm}
	P27 ¹⁰⁰¹	[S/C1 ^{sc}]
	Rb .	Rbf ^{pm}
Apoptosis	BCL-2	ced-9 ^C

Cell Cycle/DNA Damage Response Pathways The state of the s

